

# Model-independent measurement of mixing parameters in $D^0 \rightarrow K_S^0$

## $\pi^+\pi^-$ decays

LHCb Collaboration

DOI:

[10.1007/JHEP04\(2016\)033](https://doi.org/10.1007/JHEP04(2016)033)

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Document Version

Publisher's PDF, also known as Version of record

Citation for published version (Harvard):

LHCb Collaboration 2016, 'Model-independent measurement of mixing parameters in  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$  decays', *Journal of High Energy Physics*, vol. 2016, no. 33, 33. [https://doi.org/10.1007/JHEP04\(2016\)033](https://doi.org/10.1007/JHEP04(2016)033)

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RECEIVED: October 7, 2015

REVISED: February 3, 2016

ACCEPTED: March 2, 2016

PUBLISHED: April 6, 2016

# Model-independent measurement of mixing parameters in $D^0 \rightarrow K_S^0 \pi^+ \pi^-$ decays



## The LHCb collaboration

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**ABSTRACT:** The first model-independent measurement of the charm mixing parameters in the decay  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$  is reported, using a sample of  $pp$  collision data recorded by the LHCb experiment, corresponding to an integrated luminosity of  $1.0 \text{ fb}^{-1}$  at a centre-of-mass energy of 7 TeV. The measured values are

$$\begin{aligned} x &= (-0.86 \pm 0.53 \pm 0.17) \times 10^{-2}, \\ y &= (+0.03 \pm 0.46 \pm 0.13) \times 10^{-2}, \end{aligned}$$

where the first uncertainties are statistical and include small contributions due to the external input for the strong phase measured by the CLEO collaboration, and the second uncertainties are systematic.

**KEYWORDS:** Charm physics, Oscillation, Flavor physics, Hadron-Hadron scattering

**ARXIV EPRINT:** [1510.01664](https://arxiv.org/abs/1510.01664)

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## 1 Introduction

Mixing occurs in weakly decaying neutral mesons for which the flavour eigenstates of the particle and antiparticle (e.g.  $D^0$  and  $\bar{D}^0$ ) are not distinguished by any conserved quantum number. It is characterised by the differences in mass,  $\Delta M$ , and width,  $\Delta\Gamma$ , between the mass eigenstates. In the charm system these are usually expressed in a reduced form,  $x \equiv \Delta M/\Gamma$  and  $y \equiv \Delta\Gamma/(2\Gamma)$ , where  $\Gamma$  is the average of the two widths.

Mixing in charm has been observed with a significance above five standard deviations in several independent measurements [1–4] and the constraints on  $(x, y)$  are now rather precise [5]. However, most of the measurements are sensitive to  $(x^2 + y^2)$  or to  $y$  (in the limit of negligible  $CP$  violation), leading to an ambiguity in the sign of  $x$ . One approach to resolve this ambiguity is to exploit the decay to the three-body, self-conjugate final state  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$  [6–8].

The advantage of decays such as  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$  is that both Cabibbo-favoured (CF) and doubly Cabibbo-suppressed (DCS) components are present in the same final state. Therefore the strong phase differences between contributing amplitudes — and hence between mixed and unmixed decays — can be measured with an amplitude analysis [7–10] of the same data sample used to obtain the mixing parameters. This is the approach that has

been used to date. A second method, proposed in ref. [11] and building upon a related approach for determining the unitarity triangle angle  $\gamma$  [12], uses measurements of the average strong phase difference in regions of the phase space. These can be obtained from an  $e^+e^-$  collider operating at the  $\psi(3770)$  resonance. CLEO has made suitable measurements [13] and a similar study could be carried out with the larger BESIII [14]  $\psi(3770)$  sample. The advantage of this second method is that no amplitude analysis is needed: the systematic uncertainty associated with the amplitude model is replaced with the uncertainty on the strong phase measurements. It has been estimated that with BESIII data this external uncertainty should be smaller than the statistical uncertainty for  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$  yields of up to 10–20 million [15], far larger than those available today. This paper describes the first measurement of  $x$  and  $y$  with this novel method, using promptly produced charm mesons in the decay chain  $D^{*+} \rightarrow D^0 \pi^+$ ,  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$ ,  $K_S^0 \rightarrow \pi^+ \pi^-$  (charge conjugate processes are included implicitly unless otherwise noted). A sample of  $pp$  collision data recorded by the LHCb experiment in 2011 is used, corresponding to an integrated luminosity of  $1.0 \text{ fb}^{-1}$  at a centre-of-mass energy of 7 TeV.

## 2 Formalism

The formalism for the method has been presented previously [11, 13, 15], but is summarised here for clarity. The flavour eigenstates,  $|D^0\rangle$  and  $|\bar{D}^0\rangle$ , are related to the mass eigenstates,  $|D_1\rangle$  and  $|D_2\rangle$ , via

$$|D_1\rangle = p|D^0\rangle - q|\bar{D}^0\rangle, \quad (2.1)$$

$$|D_2\rangle = p|D^0\rangle + q|\bar{D}^0\rangle, \quad (2.2)$$

where  $|p|^2 + |q|^2 = 1$ . In the limit of  $CP$  conservation,  $|p/q| = 1$ . There is one free phase that is fixed by stipulating that in the limit of no indirect  $CP$  violation,  $q/p = +1$  and  $|D_1\rangle$  is the  $CP$ -odd eigenstate. The sign convention adopted for the mixing parameters is

$$x = (M_2 - M_1)/\Gamma, \quad (2.3)$$

$$y = (\Gamma_2 - \Gamma_1)/(2\Gamma). \quad (2.4)$$

For a state that is initially pure  $D^0$  at  $t = 0$ , let the state at some later time  $t$  be  $|D^0(t)\rangle$ . Likewise, let the time evolution of  $\bar{D}^0$  be  $|\bar{D}^0(t)\rangle$ . These may be evaluated as

$$|D^0(t)\rangle = g_+(t)|D^0\rangle + \frac{q}{p}g_-(t)|\bar{D}^0\rangle, \quad (2.5)$$

$$|\bar{D}^0(t)\rangle = \frac{p}{q}g_-(t)|D^0\rangle + g_+(t)|\bar{D}^0\rangle, \quad (2.6)$$

where

$$g_{\pm}(t) \equiv \frac{e^{-i(M_2 - i\Gamma_2/2)t} \pm e^{-i(M_1 - i\Gamma_1/2)t}}{2}. \quad (2.7)$$

The phase space for the three-body decay of a  $D^0$  or  $\bar{D}^0$  meson to  $K_S^0 \pi^+ \pi^-$  is conventionally represented as a Dalitz plot and can be described by two variables,  $m_{12}^2 =$

$m^2(K_S^0 \pi^+)$  and  $m_{13}^2 = m^2(K_S^0 \pi^-)$ . Let the amplitude for a  $D^0$  decay to a point  $(m_{12}^2, m_{13}^2)$  in the phase space be  $\mathcal{A}_{D^0}(m_{12}^2, m_{13}^2)$ . Neglecting direct  $CP$  violation, the amplitudes for  $D^0$  and  $\bar{D}^0$  are related by the exchange  $m_{12}^2 \leftrightarrow m_{13}^2$ ,

$$\mathcal{A}_{\bar{D}^0}(m_{12}^2, m_{13}^2) = \mathcal{A}_{D^0}(m_{13}^2, m_{12}^2). \quad (2.8)$$

In the expressions that follow, the explicit dependence of the amplitude terms  $\mathcal{A}_{D^0}$  and  $\mathcal{A}_{\bar{D}^0}$  on  $m_{12}^2$  and  $m_{13}^2$  is omitted. The amplitude  $\mathcal{A}_{D^0}(m_{12}^2, m_{13}^2, t)$  for a state that was initially  $D^0$  to decay at some later time  $t$  to a point  $(m_{12}^2, m_{13}^2)$  in the phase space is

$$\mathcal{A}_{D^0}(m_{12}^2, m_{13}^2, t) = \mathcal{A}_{D^0} g_+(t) + \frac{q}{p} \mathcal{A}_{\bar{D}^0} g_-(t). \quad (2.9)$$

Similarly,

$$\mathcal{A}_{\bar{D}^0}(m_{12}^2, m_{13}^2, t) = \mathcal{A}_{\bar{D}^0} g_+(t) + \frac{p}{q} \mathcal{A}_{D^0} g_-(t). \quad (2.10)$$

The probability density  $\mathcal{P}_{D^0}(m_{12}^2, m_{13}^2, t)$  is given by the modulus squared of the amplitude multiplied by a normalisation factor of  $\Gamma$ ,

$$\mathcal{P}_{D^0}(m_{12}^2, m_{13}^2, t) = \Gamma |\mathcal{A}_{D^0}(m_{12}^2, m_{13}^2, t)|^2, \quad (2.11)$$

with  $\mathcal{P}_{\bar{D}^0}$  defined similarly in terms of  $\mathcal{A}_{\bar{D}^0}$ . Performing a Taylor expansion and neglecting terms of order  $x^2$ ,  $xy$ , and  $y^2$ , these evaluate to

$$\mathcal{P}_{D^0}(m_{12}^2, m_{13}^2, t) = \Gamma e^{-\Gamma t} \left[ |\mathcal{A}_{D^0}|^2 - \Gamma t \operatorname{Re} \left( \frac{q}{p} \mathcal{A}_{D^0}^* \mathcal{A}_{\bar{D}^0}(y + ix) \right) \right], \quad (2.12)$$

$$\mathcal{P}_{\bar{D}^0}(m_{12}^2, m_{13}^2, t) = \Gamma e^{-\Gamma t} \left[ |\mathcal{A}_{\bar{D}^0}|^2 - \Gamma t \operatorname{Re} \left( \frac{p}{q} \mathcal{A}_{D^0} \mathcal{A}_{\bar{D}^0}^*(y + ix) \right) \right]. \quad (2.13)$$

Neglecting  $CP$  violation for the purposes of the mixing measurement,  $q/p = 1$  and hence

$$\mathcal{P}_{D^0}(m_{12}^2, m_{13}^2, t) = \Gamma e^{-\Gamma t} \left[ |\mathcal{A}_{D^0}|^2 - \Gamma t \operatorname{Re} (\mathcal{A}_{D^0}^* \mathcal{A}_{\bar{D}^0}(y + ix)) \right], \quad (2.14)$$

$$\mathcal{P}_{\bar{D}^0}(m_{12}^2, m_{13}^2, t) = \Gamma e^{-\Gamma t} \left[ |\mathcal{A}_{\bar{D}^0}|^2 - \Gamma t \operatorname{Re} (\mathcal{A}_{\bar{D}^0} \mathcal{A}_{D^0}^*(y + ix)) \right]. \quad (2.15)$$

These densities may be integrated over regions of the phase space. Various binning schemes are possible; this analysis uses the one referred to as the “equal  $\Delta\delta_D$  BaBar 2008” binning in ref. [13], in which the strong phase variation within each bin of the phase space is minimised. This has the advantage of reducing the sensitivity to detector effects such as variation in efficiency across the phase space. In this scheme there are 16 bins, with bins 1 to 8 in the region of the phase space  $m_{12}^2 > m_{13}^2$  and bins  $-1$  to  $-8$  in the region  $m_{12}^2 < m_{13}^2$ . The bins are symmetric about the leading diagonal, with bin  $i$  mapped to bin  $-i$  by the transformation  $(m_{12}^2, m_{13}^2) \rightarrow (m_{13}^2, m_{12}^2)$ . The quantities  $T_i$  and  $X_i$  are defined by the integrals

$$T_i \equiv \int_i |\mathcal{A}_{D^0}|^2 dm_{12}^2 dm_{13}^2, \quad (2.16)$$

$$X_i \equiv \frac{1}{\sqrt{T_i T_{-i}}} \int_i \mathcal{A}_{D^0}^* \mathcal{A}_{\bar{D}^0} dm_{12}^2 dm_{13}^2, \quad (2.17)$$

and the  $X_i$  may in turn be expressed in terms of real quantities  $c_i$  and  $s_i$  as

$$c_i \equiv \text{Re}(X_i), \quad (2.18)$$

$$s_i \equiv -\text{Im}(X_i). \quad (2.19)$$

Given the symmetric binning, eq. (2.8) implies that  $X_{-i} = X_i^*$ , and thus  $c_{-i} = c_i$  and  $s_{-i} = -s_i$ .

With these definitions, the integrated probability densities are

$$\begin{aligned} \mathcal{P}_{D^0}(i; t) &= \int_i \mathcal{P}_{D^0}(m_{12}^2, m_{13}^2, t) dm_{12}^2 dm_{13}^2 \\ &= \Gamma e^{-\Gamma t} \left[ T_i - \Gamma t \sqrt{T_i T_{-i}} \{y c_i + x s_i\} \right], \end{aligned} \quad (2.20)$$

and

$$\mathcal{P}_{\bar{D}^0}(i; t) = \Gamma e^{-\Gamma t} \left[ T_{-i} - \Gamma t \sqrt{T_i T_{-i}} \{y c_i - x s_i\} \right]. \quad (2.21)$$

These distributions are used to obtain the mixing parameters  $x$  and  $y$ . The values of  $T_i$ ,  $c_i$ , and  $s_i$  measured by the CLEO collaboration are given in tables VII and XVI of ref. [13].<sup>1</sup>

### 3 Detector, selection and simulation

The LHCb detector [16, 17] is a single-arm forward spectrometer covering the pseudo-rapidity range  $2 < \eta < 5$ , designed for the study of particles containing  $b$  or  $c$  quarks. The detector includes a high-precision tracking system consisting of a silicon-strip vertex detector surrounding the  $pp$  interaction region, a large-area silicon-strip detector located upstream of a dipole magnet with a bending power of about 4 Tm, and three stations of silicon-strip detectors and straw drift tubes placed downstream of the magnet. The tracking system provides a measurement of momentum,  $p$ , of charged particles with a relative uncertainty that varies from 0.5% at low momentum to 1.0% at 200 GeV/ $c$ . The minimum distance of a track to a primary vertex, the impact parameter (IP), is measured with a resolution of  $(15 + 29/p_T) \mu\text{m}$ , where  $p_T$  is the component of the momentum transverse to the beam, in GeV/ $c$ . Different types of charged hadrons are distinguished using information from two ring-imaging Cherenkov detectors. Photons, electrons and hadrons are identified by a calorimeter system consisting of scintillating-pad and preshower detectors, an electromagnetic calorimeter and a hadronic calorimeter. Muons are identified by a system composed of alternating layers of iron and multiwire proportional chambers.

The online event selection is performed by a trigger [18], which consists of a hardware stage, based on information from the calorimeter and muon systems, followed by a software stage, which applies a full event reconstruction. At the hardware trigger stage, events are required to have a muon with high  $p_T$  or a hadron, photon or electron with high transverse energy in the calorimeters. In the subsequent software trigger, pairs of oppositely charged tracks are combined to form  $K_S^0$  candidates, and those are in turn combined with a second pair of oppositely charged tracks to form  $D^0$  candidates. For the 2011 dataset, the trigger

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<sup>1</sup>Note that the captions for tables VII and VIII were exchanged in ref. [13], and that the supplementary material defining the binning contains an off-by-one error in the bin indices.

requires that all four tracks be reconstructed in the vertex detector, reducing the  $K_s^0$  efficiency significantly. Both the  $K_s^0$  and  $D^0$  candidate vertices are required to be displaced from any primary  $pp$  interaction vertex (PV) in the event, and additional geometrical and kinematic criteria are applied to suppress background and ensure consistency with a  $D^0 \rightarrow K_s^0 \pi^+ \pi^-$  decay. These include a requirement that at least one of the four tracks has an impact parameter larger than  $100 \mu\text{m}$  with respect to any PV.

After offline processing, additional selection criteria are applied to further suppress background. These include particle identification requirements on the  $D^0$  daughter tracks, as well as requirements that the track and vertex fits be of good quality, that the  $K_s^0$  vertex be at least 10 mm downstream of the  $D^0$  vertex, that the  $K_s^0$  candidate mass lie within  $\pm 11.4 \text{ MeV}/c^2$  of the known value [19], that the  $D^0$  candidate mass  $m_D$  lie within  $\pm 85 \text{ MeV}/c^2$  of the known value [19], and that the reconstructed  $D^0$  decay time  $t_D$  lie within  $0.3 < t_D < 5 \text{ ps}$ . The  $D^0$  candidate is also required to have no more than two turning points in its decay time acceptance function (see section 4.3). It is then combined with a fifth pion track, referred to as the soft pion, to form a  $D^{*+}$  candidate. Both the soft pion and  $D^0$  candidate are constrained to originate from the same PV. Good vertex fit quality is required, and particle identification requirements are applied to the soft pion. The mass difference  $\Delta m = m_{D^{*+}} - m_D$  is required to lie within the range  $m_\pi < \Delta m < (m_\pi + 15 \text{ MeV}/c^2)$ , where  $m_{D^{*+}}$  is the mass of the  $D^{*+}$  candidate and  $m_\pi$  is the charged pion mass. If there is more than one distinct  $D^0 \rightarrow K_s^0 \pi^+ \pi^-$  candidate then one is chosen at random and the rest are discarded. If, after this, there are multiple  $D^{*\pm}$  candidates then the one with the best vertex fit quality is retained and the rest are discarded.

Simulated events are used for cross-checks. In the simulation,  $pp$  collisions are generated using PYTHIA 6 [20] with a specific LHCb configuration [21]. Decays of hadronic particles are described by EVTGEN [22], in which final-state radiation is generated using PHOTOS [23]. The interaction of the generated particles with the detector, and its response, are implemented using the GEANT4 toolkit [24, 25] as described in ref. [26].

## 4 Fits

### 4.1 Overview

The mixing parameters  $x$  and  $y$  are determined by a sequence of fits to the distributions of the variables  $(m_D, \Delta m)$  and  $(t_D, \ln \chi_{\text{IP}}^2)$ , initially in the whole phase space and later in individual regions. The impact parameter  $\chi^2$ ,  $\chi_{\text{IP}}^2$ , is defined as the difference in the vertex fit  $\chi^2$  of the associated PV with and without the  $D^0$  candidate. It is used to separate prompt charm that originates at the PV from secondary charm produced at a displaced vertex. The dominant source of secondary charm is from decays of  $b$ -hadrons. Two other variables are also used to describe the per-event decay time acceptance function, introduced in section 4.3. Unless otherwise specified, all data passing the selection described in section 3 are used. Where reference is made to a narrow signal window in  $m_D$  or  $\Delta m$ , this corresponds to a stricter requirement:  $\pm 20 \text{ MeV}/c^2$  around the known  $D^0$  mass, or  $144.2 <$

$\Delta m < 146.4 \text{ MeV}/c^2$ . The mass sidebands are defined as  $1785 < m_D < 1810 \text{ MeV}/c^2$  and  $1920 < m_D < 1945 \text{ MeV}/c^2$ .

First, an extended maximum likelihood fit to the  $m_D$  distribution of all selected  $D^{*+}$  candidates is performed to determine the amounts of  $D^0$  signal and combinatorial background in the narrow  $m_D$  signal window (section 4.2). Second is a maximum likelihood fit to the  $(t_D, \ln \chi_{\text{IP}}^2)$  distribution of those candidates in the narrow  $m_D$  signal window, using the mass sidebands to estimate the background distributions (section 4.4). This fit uses the yields determined in the first fit, and serves to determine the  $\ln \chi_{\text{IP}}^2$  shapes for prompt and secondary charm. It is not sensitive to mixing. Third is a set of 32 extended maximum likelihood fits, each to the  $(m_D, \Delta m)$  distribution in a particular phase space bin, with the  $D^{*+}$  and  $D^{*-}$  samples fitted separately (section 4.5). Each fit provides measurements of the amounts of signal and background in the narrow  $(m_D, \Delta m)$  window for the corresponding bin. Fourth is a simultaneous maximum likelihood fit to the  $(t_D, \ln \chi_{\text{IP}}^2)$  distributions of candidates for the 32 subsamples (section 4.6). Signal candidates are required to lie in the narrow  $m_D$  and  $\Delta m$  signal windows, with the mass sidebands used to constrain the combinatorial background. This fit uses the  $\ln \chi_{\text{IP}}^2$  shapes from the second fit and the yield estimates from the third fit, and produces measurements of the mixing parameters  $x$  and  $y$ .

Only the fit procedure and results are discussed in this section. Cross-checks and systematic uncertainties are discussed in section 5. All aspects of the selection and fit procedure were finalised before any measurements of  $x$  and  $y$  were made. Unless otherwise stated, all parameters introduced are left free in the fits.

## 4.2 Fit to $m_D$

The probability density functions (PDFs) used to model the  $m_D$  distributions are expressed in terms of exponential, Gaussian ( $G$ ), bifurcated Gaussian ( $B$ ), and Crystal Ball ( $C$ ) [27] functions. Only two components are needed:  $D^0$  signal and combinatorial background. The PDF for  $D^0$  signal (sig) is the sum of a Gaussian, a bifurcated Gaussian, and a Crystal Ball function,

$$f_1(m_D|\text{sig}) = \eta_1 G(m_D; \mu_D, \sigma_1) + \eta_2 B(m_D; \mu_D, \sigma_L, \sigma_R) + (1 - \eta_1 - \eta_2) C(m_D; \mu_D, \sigma_2, \alpha, n), \quad (4.1)$$

where the order of the Crystal Ball function,  $n$ , is fixed to three. The PDF for the combinatorial background,  $f_1(m_D|\text{cmb})$ , is an exponential function. The total PDF is then

$$f_1(m_D) = P_1(\text{sig}) f_1(m_D|\text{sig}) + P_1(\text{cmb}) f_1(m_D|\text{cmb}), \quad (4.2)$$

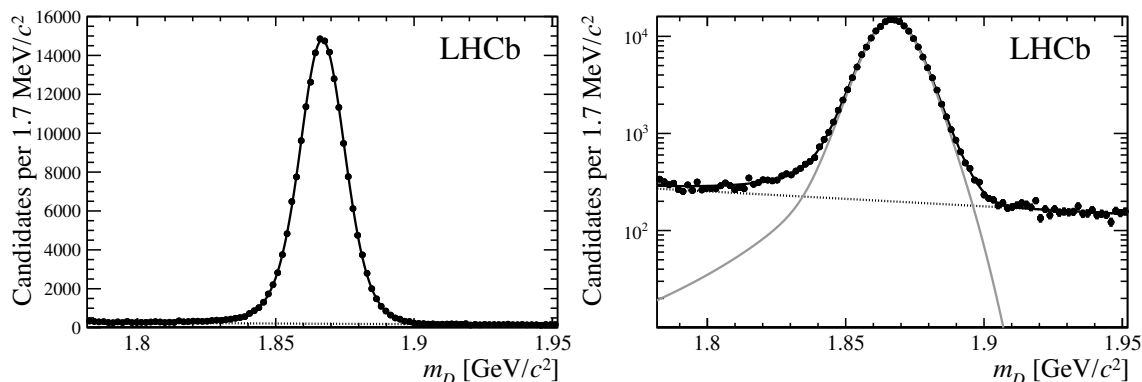
where  $P_1(\text{sig})$  and  $P_1(\text{cmb})$  describe the fractions of signal and background in the data sample used for the first fit, and sum to unity.

The results of the first fit are shown in figure 1. The fit yields 178k signal events within the narrow  $m_D$  signal window, and the purity within this window is  $(97.4 \pm 0.3)\%$ .

## 4.3 Time acceptance correction

The probability for a  $D^{*+}$  signal decay to be successfully triggered, reconstructed, and selected depends upon the decay time of its  $D^0$  daughter. The time-dependent fits must,





**Figure 1.** Fitted  $m_D$  distribution. Both plots show the same data sample with (left) linear and (right) logarithmic vertical scales. The curves show the results of the first fit, described in section 4.2: the total (solid black), the background component (dotted), and the signal component (grey, right only).

therefore, take account of the nonuniform decay time acceptance. A data-driven method referred to as swimming [28] is used. This approach follows that used in previous LHCb measurements of the mixing and indirect  $CP$  violation parameters,  $y_{CP}$  and  $A_\Gamma$ , in  $D^0$  decays [29, 30], and at previous experiments [31–34].

The principle of the method is that the decay time acceptance is determined by selection criteria that can be reproduced later. The criteria for this analysis are given in section 3. (In practice they are applied to the measured rather than the true decay time; the resolution is neglected and considered as a systematic effect (section 5).) Those criteria can be tested again after modifying the candidate — specifically, with a different decay time. By repeatedly testing the criteria for many decay time values spanning the allowed range, the acceptance function for an individual candidate may be determined empirically. Aside from a correction factor discussed later in this section, the value of this function is 1 for those decay times at which all of the criteria are fulfilled, and 0 at all other times. Since candidates with  $t_D < 0.3$  ps are rejected, the acceptance function is zero below that point. It must also be zero at very large decay times, both because of the upper bound on  $t_D$  and because of the finite length of the vertex detector. Therefore, the acceptance function will take the form of a top-hat function  $[\Theta(t_D - t_0) - \Theta(t_D - t_1)]$ , where  $\Theta$  is the Heaviside function and  $t_1 > t_0$ , or will be the sum of several nonoverlapping top-hat functions. The decay times at which the acceptance changes between 0 and 1 are referred to as the turning points.

For approximately 90% of selected candidates, the acceptance is a single top-hat with exactly two turning points. The remaining candidates have a more complicated acceptance function, typically due to the presence of a second  $pp$  primary vertex nearby. As in the previous analysis using this technique [30], candidates with more than two turning points are rejected. This enables a more robust description of the turning point variable distributions (see below) and suppresses events in which the primary vertex association is ambiguous.

The implementation of the decay time acceptance calculation is simplified by a number of assumptions. First, the hardware triggers do not depend on the  $D^0$  decay time and can

therefore be ignored when evaluating the acceptance function. Second, the decay time acceptance depends only on the  $D^0$  reconstruction and selection: it is not affected by the soft pion and  $D^{*+}$  requirements. Third, the full vertex detector pattern recognition is not re-run when changing the  $D^0$  decay time; instead, the changes to the decay geometry are made analytically. Requirements on the number of hits on a track in the vertex detector subsystem are approximated as requirements that the modified trajectory pass through a corresponding number of subdetector modules. Finally, an additional correction factor  $\varepsilon(t_D)$  is applied to the acceptance function to model the effect of a track quality cut in the reconstruction, which reduces the efficiency for tracks produced further from the beam axis. The correction is derived from samples of simulated events and is parameterised as a polynomial function.

For an individual event, the acceptance function can be written as

$$a(t_D; t_0, \Delta t) = [\Theta(t_D - t_0) - \Theta(t_D - t_0 - \Delta t)] \varepsilon(t_D), \quad (4.3)$$

where  $t_0$  is the first turning point (TP) and  $\Delta t$  is the difference between the two turning points. Although the acceptance function is determined for each event independently, models of the distribution  $f_{\text{TP}}(t_0, \Delta t)$  of the turning point variables  $t_0$  and  $\Delta t$  are required for the decay time fits. The distribution is assumed to factorise,

$$f_{\text{TP}}(t_0, \Delta t) = f_{\text{TP},0}(t_0) f_{\text{TP},\Delta}(\Delta t). \quad (4.4)$$

Nonparametric functions are used to model the turning point PDFs. The distribution  $f_{\text{TP},0}(t_0)$  is modelled as a histogram PDF with 100 bins spanning the range 0–3 ps and the distribution  $f_{\text{TP},\Delta}(\Delta t)$  is modelled as a one-dimensional Gaussian kernel PDF [35]. The same method is used for all components, and is based on data in the mass sidebands for combinatorial background. Candidates in the narrow mass signal window are used for prompt and secondary  $D^0$  mesons, both of which are assumed to have the same turning point distribution in the baseline fit.

#### 4.4 Separation of prompt and secondary candidates

The second fit is used to determine the relative proportions of prompt and secondary  $D^0$  signal, and to model their  $\ln \chi^2_{\text{TP}}$  distributions. It also serves as an important cross-check since it allows the mean  $D^0$  lifetime to be computed in the  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$  sample. No distinction is made in the fit between  $D^0$  and  $\bar{D}^0$  candidates, nor between different regions of the phase space, so by design it is insensitive to mixing. While a dominance of  $CP$ -odd or of  $CP$ -even components in the final state could in principle shift the mean lifetime by up to  $\pm y \Gamma \approx 2.5$  fs, the net  $CP$  has recently been shown to be almost zero [36] so that the effective lifetime is close to  $\tau_D$ . Similarly, previous amplitude analyses all found that the decay is dominated by flavour-specific processes, with total fit fractions of about 70% [7–10], implying that the maximum scale of the effect is below the sensitivity of this analysis.

In this fit, the underlying decay time distribution for the prompt (prm)  $D^0$  signal is taken to be an exponential function for  $t_D > 0$  with characteristic time  $\tau_D$ . For a

particular event, the expected  $t_D$  distribution is this exponential multiplied by the per-event acceptance function given in eq. (4.3),

$$f_2(t_D|t_0, \Delta t; \text{prm}) = n a(t_D; t_0, \Delta t) e^{-t_D/\tau_D}, \quad (4.5)$$

where  $n$  is a normalisation factor and the decay time resolution has been neglected. Note that the expression in eq. (4.5) depends explicitly on the turning point variables  $t_0$  and  $\Delta t$ . To separate out this dependence, the models for the turning point distributions given in section 4.3 are used. The PDF for prompt charm may then be written as

$$f_2(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2 | \text{prm}) = f_2(\ln \chi_{\text{IP}}^2 | t_D; \text{prm}) f_2(t_D | t_0, \Delta t; \text{prm}) \times f_{\text{TP},0}(t_0 | D) f_{\text{TP},\Delta}(\Delta t | D), \quad (4.6)$$

where  $D$  denotes PDFs used for both prompt and secondary  $D^0$ , and  $f_2(\ln \chi_{\text{IP}}^2 | t_D; \text{prm})$  is a parameterisation of the  $\ln \chi_{\text{IP}}^2$  distribution for a given decay time, taking the form

$$f_2(\ln \chi_{\text{IP}}^2 | t_D; \text{prm}) = \eta G(\ln \chi_{\text{IP}}^2; \mu_p(t_D), \sigma_1) + (1 - \eta) B(\ln \chi_{\text{IP}}^2; \mu_p(t_D), \sigma_L, \sigma_R), \quad (4.7)$$

where  $\mu_p(t_D)$ , the most probable value of  $\ln \chi_{\text{IP}}^2$ , is a linear function.

A similar approach is used for the secondary (sec)  $D^0$  signal, except that the underlying decay time distribution is taken to be the convolution of two exponential functions restricted to  $t_D > 0$  and with characteristic times  $\tau_1$  and  $\tau_2$ . Since  $[\Theta(t_D) e^{-t_D/\tau_1}] \otimes [\Theta(t_D) e^{-t_D/\tau_2}]$  may be rewritten as  $(e^{-t_D/\tau_2} - e^{-t_D/\tau_1})$  with an appropriate normalisation factor, the expression remains analytically integrable and takes the form

$$f_2(t_D | t_0, \Delta t; \text{sec}) = n a(t_D; t_0, \Delta t) \left( e^{-t_D/\tau_2} - e^{-t_D/\tau_1} \right), \quad (4.8)$$

where  $n$  is again a normalisation factor. The  $\ln \chi_{\text{IP}}^2$  distribution also differs from that used for prompt charm,

$$f_2(\ln \chi_{\text{IP}}^2 | t_D; \text{sec}) = \eta G(\ln \chi_{\text{IP}}^2; \mu_s(t_D), \alpha \sigma_1) + (1 - \eta) B(\ln \chi_{\text{IP}}^2; \mu_s(t_D), \alpha (\sigma_L + \beta t_D), \alpha \sigma_R). \quad (4.9)$$

Compared to eq. (4.7), the width of the peak is multiplied by  $\alpha$ , with the lower tail of the bifurcated Gaussian having a further, time-dependent broadening. In addition, the decay time at which the function is maximised,  $\mu_s(t_D)$ , is taken empirically to evolve as

$$\mu_s(t_D) = \mu_{s0} + B(1 - e^{C t_D}). \quad (4.10)$$

Using the models for the turning point distributions given in section 4.3, the PDF for secondary charm may be written as

$$f_2(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2 | \text{sec}) = f_2(\ln \chi_{\text{IP}}^2 | t_D; \text{sec}) f_2(t_D | t_0, \Delta t; \text{sec}) \times f_{\text{TP},0}(t_0 | D) f_{\text{TP},\Delta}(\Delta t | D). \quad (4.11)$$

The combinatorial background is described in a different way. To begin, a nonparametric distribution is fitted to the data in the mass sidebands. However, this model, a two-dimensional Gaussian kernel function, cannot be used directly in the fit: the PDF

used must depend explicitly on the turning point variables [37]. Therefore, an unfolding procedure is applied to obtain the underlying decay time distribution before acceptance effects. The acceptance is then incorporated in the same way as for the other components. The PDF for combinatorial background may be written as

$$f_2(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2 | \text{cmb}) = f_2(\ln \chi_{\text{IP}}^2 | t_D; \text{cmb}) f_2(t_D | t_0, \Delta t; \text{cmb}) \times f_{\text{TP},0}(t_0 | \text{cmb}) f_{\text{TP},\Delta}(\Delta t | \text{cmb}), \quad (4.12)$$

where  $f_{\text{TP},0}(t_0 | \text{cmb})$  and  $f_{\text{TP},\Delta}(\Delta t | \text{cmb})$  are obtained as described in section 4.3, and  $f_2(\ln \chi_{\text{IP}}^2 | t_D; \text{cmb})$  and  $f_2(t_D | t_0, \Delta t; \text{cmb})$  are derived from the distributions in the mass sidebands as described above.

Combining the above, the total PDF used in the fit is

$$f_2(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2) = \sum_j f_2(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2 | j) P_2(j), \quad (4.13)$$

where the index  $j$  runs over the prompt, secondary, and combinatoric components and  $\sum_j P_2(j) = 1$ . The value of  $P_2(\text{cmb})$  is fixed based on the results of the preceding fit to  $m_D$ . The sum  $[P_2(\text{prm}) + P_2(\text{sec})]$  is likewise fixed, but with the secondary fraction of the signal free.

Pseudoexperiments are used to validate the fit procedure. In each pseudoexperiment, events from each category (prompt  $D^0$  mesons, secondary  $D^0$  mesons, and combinatorial background) are generated according to the expected distributions and analysed following the same procedure as used for data, including estimation of the per-event decay time acceptance function with the swimming method. In an ensemble of approximately 500 pseudoexperiments generated assuming a true  $D^0$  lifetime of 410 ps, the mean of the fitted values of  $\tau_D$  is  $409.92 \pm 0.06$  fs, and the normalised residuals are described by a Gaussian distribution with a mean of  $0.016 \pm 0.049$  and a width of  $1.03 \pm 0.04$ .

Applying the fit to the data, the measured lifetime is  $\tau_D = 410.9 \pm 1.1$  fs, where the uncertainty is purely statistical. This is consistent with the world average value of  $410.1 \pm 1.5$  fs [19]. The agreement between the fit and data is shown in figure 2. An excess is seen at very long decay times, likely due to imperfect modelling of the secondary component, but there is no effect on the measurement of the lifetime of the prompt component.

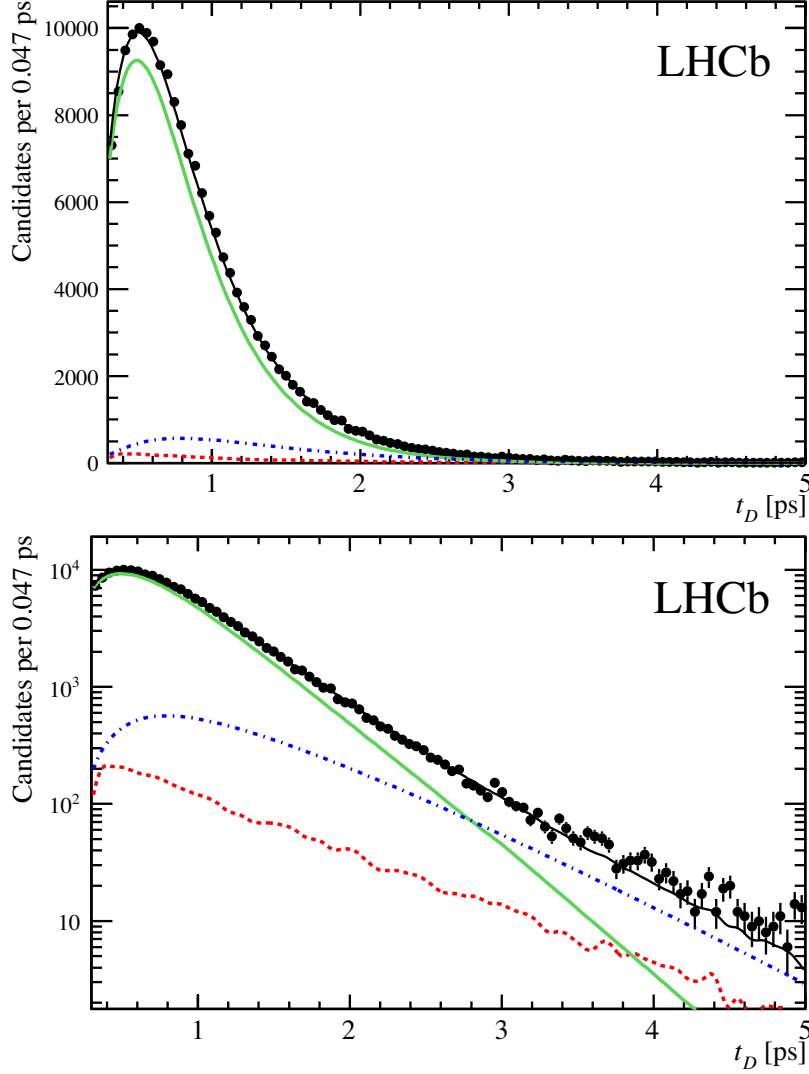
#### 4.5 Fits to $m_D$ and $\Delta m$

The third step consists of separate fits to the  $(m_D, \Delta m)$  distributions of the phase space bins. The fits include three components:  $D^{*+}$  signal (sig), background from genuine  $D^0$  that are combined with an unrelated soft pion (Dbg), and combinatorial background (cmb). In each case, the PDF is assumed to factorise into  $m_D$ -dependent and  $\Delta m$ -dependent terms. The three components may be written as

$$f_3(m_D, \Delta m | \text{sig}) = f_3(m_D | \text{peak}) f_3(\Delta m | \text{peak}), \quad (4.14)$$

$$f_3(m_D, \Delta m | \text{Dbg}) = f_3(m_D | \text{peak}) f_3(\Delta m | \text{smooth}), \quad (4.15)$$

$$f_3(m_D, \Delta m | \text{cmb}) = f_3(m_D | \text{smooth}) f_3(\Delta m | \text{smooth}), \quad (4.16)$$



**Figure 2.** Decay time projection from the fit for separation of prompt and secondary candidates. The curves show the results of the fit described in section 4.4: the total (solid black), the prompt component (solid green), the secondary component (dot-dashed blue), and the combinatorial component (dashed red). Both plots show the same data sample with linear (top) and logarithmic (bottom) vertical scales.

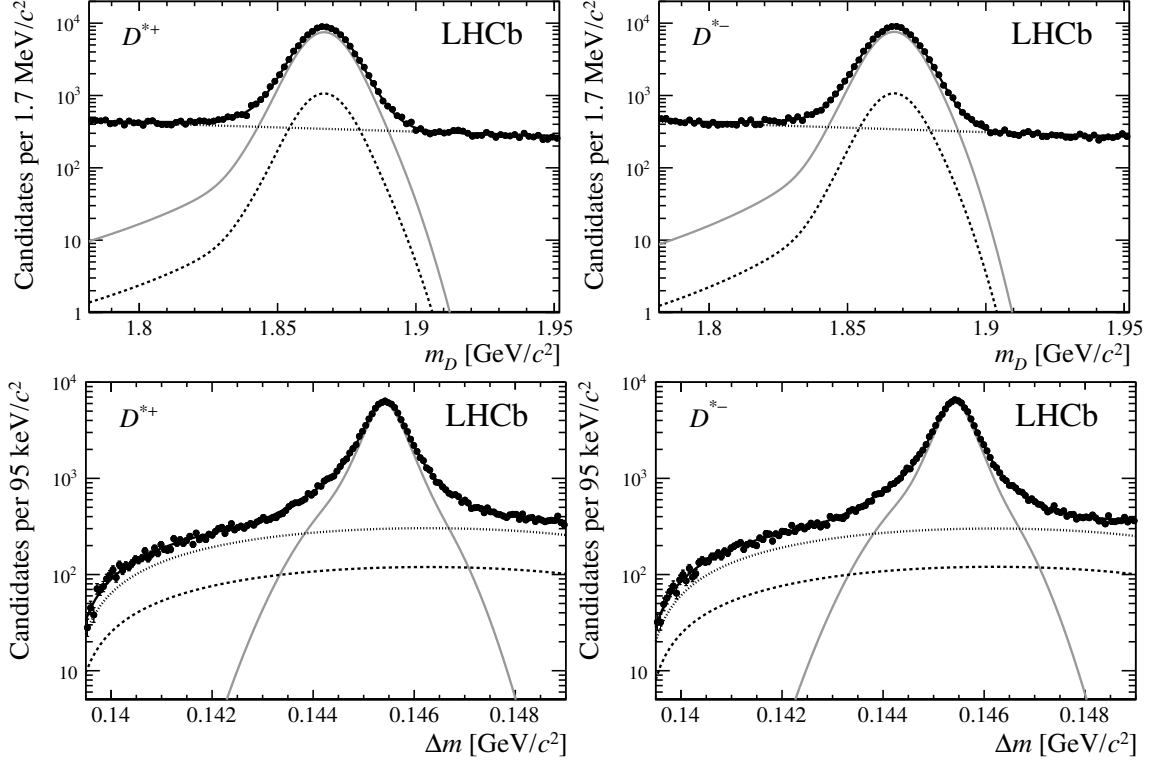
where the peaking components are defined as

$$f_3(m_D|\text{peak}) = \eta_1 G(m_D; \mu_D, \sigma_1) + \eta_2 G(m_D; \mu_D, \sigma_2) + (1 - \eta_1 - \eta_2) C(m_D; \mu_D, \sigma_3, \alpha, n), \quad (4.17)$$

$$f_3(\Delta m|\text{peak}) = \eta_3 G(\Delta m; \mu_{\Delta m}, \sigma_4) + \eta_4 G(\Delta m; \mu_{\Delta m}, \sigma_5) + (1 - \eta_3 - \eta_4) B(\Delta m; \mu_{\Delta m}, \sigma_L, \sigma_R). \quad (4.18)$$

For the nonpeaking components,  $f_3(m_D|\text{smooth})$  is an exponential function and  $f_3(\Delta m|\text{smooth})$  is a second-order polynomial. The total PDF may then be written as

$$f_3(m_D, \Delta m) = \sum_j f_3(m_D, \Delta m|j) P_3(j), \quad (4.19)$$



**Figure 3.** Fitted  $(m_D, \Delta m)$  distributions. The upper row shows the  $m_D$  projection and the lower row  $\Delta m$ . The left column shows  $D^{*+}$  candidates and the right column  $D^{*-}$ . The signal and background components are shown separately (signal as solid grey,  $D^0$  background dashed, combinatoric background dotted, and the sum as solid black).

where the index  $j$  runs over the signal,  $D^0$  background, and combinatoric components, and  $\sum_j P_3(j) = 1$ .

To avoid an excessive number of free parameters when splitting the data into many independent subsamples, the third fit is done in two stages. Initially, fits to  $f_3(m_D, \Delta m)$  are done without dividing the data by phase space bin such that there are only two subsamples,  $D^{*+}$  and  $D^{*-}$ . The results of these fits are shown in figure 3 and correspond to yields of approximately 85k each of  $D^{*+}$  and  $D^{*-}$  within the narrow signal window. The parameters for  $f_3(m_D|\text{peak})$ ,  $f_3(\Delta m|\text{peak})$ , and  $f_3(\Delta m|\text{smooth})$  are then fixed. Individual fits to each of the 32 subsamples are then carried out, with only the parameters of the combinatorial background shape,  $f_3(m_D|\text{smooth})$ , and the yield fractions  $P_3(j)$  free.

#### 4.6 Mixing parameters

The fourth fit uses the  $(t_D, \ln \chi_{\text{IP}}^2)$  distributions in each of the phase space bins for  $D^0$  and  $\bar{D}^0$  to determine the mixing parameters  $x$  and  $y$ . For a particular phase space bin  $i$  and  $D^{*\pm}$  charge  $q$ , the total PDF is

$$f_4(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2, i, q) = \sum_j f_4(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2, i, q|j) P_4(i, q, j), \quad (4.20)$$

where  $\sum_j P_4(i, q, j) = 1$  and the index  $j$  runs over the components: prompt  $D^{*\pm}$  (p-sig), prompt  $D^0$  background (p-Dbg), secondary  $D^{*\pm}$  (s-sig), secondary  $D^0$  background (s-Dbg), and combinatorial background (cmb).

The prompt  $D^{*\pm}$  component comprises prompt  $D^0$  or  $\bar{D}^0$  mesons whose initial flavour is correctly identified. Its underlying decay time distribution is given by  $\mathcal{P}_{D^0}(i; t_D)$  in eq. (2.20) for  $D^{*+}$  and by  $\mathcal{P}_{\bar{D}^0}(i; t_D)$  in eq. (2.21) for  $D^{*-}$ , denoted  $\mathcal{P}_q$ . Taking the time-dependent acceptance into account in the same way as was done for the second fit in eq. (4.5), the per-candidate decay time PDF is

$$f_4(t_D|t_0, \Delta t, i, q; \text{p-sig}) = n a(t_D; t_0, \Delta t) \mathcal{P}_q(i; t_D), \quad (4.21)$$

where  $n$  is a normalisation constant. The  $\ln \chi_{\text{IP}}^2$  distribution for prompt  $D^{*+}$  signal at a given decay time is fixed to that obtained in the second fit (see eq. (4.7)), as is that for prompt  $D^0$  background,

$$f_4(\ln \chi_{\text{IP}}^2|t_D; \text{p-sig}) = f_4(\ln \chi_{\text{IP}}^2|t_D; \text{p-Dbg}) = f_2(\ln \chi_{\text{IP}}^2|t_D; \text{prm}). \quad (4.22)$$

The non-parametric turning point distributions,  $f_{\text{TP},0}(t_0|i; D)$  and  $f_{\text{TP},\Delta}(\Delta t|i; D)$ , are obtained in the same way as was done for the second fit, except that each phase space bin is now considered separately; here the label  $D$  denotes that the distributions are used for all components that contain a real  $D^0$  or  $\bar{D}^0$  (p-sig, p-Dbg, s-sig, s-Dbg). The prompt  $D^{*+}$  PDF is

$$f_4(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2, i, q|\text{p-sig}) = f_4(\ln \chi_{\text{IP}}^2|t_D; \text{p-sig}) f_4(t_D|t_0, \Delta t, i, q; \text{p-sig}) \times f_{\text{TP},0}(t_0|i; D) f_{\text{TP},\Delta}(\Delta t|i; D). \quad (4.23)$$

The prompt  $D^0$  background component consists of correctly reconstructed prompt  $D^0$  (or  $\bar{D}^0$ ) mesons, each of which is paired with an unrelated soft pion such that the assigned initial flavour is random. Ignoring the assigned flavour, the underlying decay time distribution for phase space bin  $i$ ,  $u(t_D; i)$ , is a linear combination of  $\mathcal{P}_{D^0}(i; t_D)$  and  $\mathcal{P}_{\bar{D}^0}(i; t_D)$ . The coefficients depend on the relative populations of bin  $i$  for  $D^0$  and bin  $-i$  for  $\bar{D}^0$ ,  $T_i$  and  $T_{-i}$  defined in eq. (2.16), since the  $\bar{D}^0$  Dalitz plot is the mirror reflection of that of  $D^0$  neglecting  $CP$  violation. The underlying decay time distribution is thus

$$u(t_D; i) = \frac{p_{D^0} T_i \mathcal{P}_{D^0}(i; t_D) + (1 - p_{D^0}) T_{-i} \mathcal{P}_{\bar{D}^0}(i; t_D)}{p_{D^0} T_i + (1 - p_{D^0}) T_{-i}}, \quad (4.24)$$

where  $p_{D^0}$  is the fraction of the prompt  $D^0$  background due to  $D^0$  mesons and  $(1 - p_{D^0})$  is the fraction due to  $\bar{D}^0$  mesons. Since production and detection charge asymmetries for pions in the relevant kinematic region are small [38],  $p_{D^0}$  is assumed to be 0.5. The per-candidate decay time PDF is then

$$f_4(t_D|t_0, \Delta t, i, q; \text{p-Dbg}) = n a(t_D; t_0, \Delta t) u(t_D; i), \quad (4.25)$$

where  $n$  is again a calculable normalisation factor. The turning point distributions  $f_{\text{TP},0}(t_0|\text{prompt})$  and  $f_{\text{TP},\Delta}(\Delta t|\text{prompt})$  are fixed to be the same as those obtained in

the second fit. The prompt  $D^0$  background PDF is

$$f_4(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2, i, q | \text{p-Dbg}) = f_4(\ln \chi_{\text{IP}}^2 | t_D; \text{p-Dbg}) f_4(t_D | t_0, \Delta t, i, q; \text{p-Dbg}) \times f_{\text{TP},0}(t_0 | i; D) f_{\text{TP},\Delta}(\Delta t | i; D). \quad (4.26)$$

For the secondary  $D^{*\pm}$  and secondary  $D^0$  background components, the effect of mixing is neglected so that the underlying time distribution does not depend on the identified flavour or on the phase space bin. The same functional form is used as for the second fit, and the parameters are fixed to those obtained in the second fit. Thus, the PDF is the same as that given in eq. (4.11),

$$f_4(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2, i, q | \text{s-sig}) = f_4(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2, i, q | \text{s-Dbg}) = f_2(t_0, \Delta t, t_D, \ln \chi_{\text{IP}}^2 | \text{sec}). \quad (4.27)$$

It is assumed that the fraction of  $D^{*\pm}$  signal that is from secondary production is the same in every phase space bin, and that the same fraction also applies to the secondary  $D^0$  background.

For the combinatorial component, nonparametric models are used for the decay time and  $\ln \chi_{\text{IP}}^2$  distributions in a similar way to the second fit. However, the distributions for each of the 32 subsamples, split by phase space bin and by  $D^{*\pm}$  charge, are modelled independently according to the mass sidebands for that bin and charge.

Thus, nearly all of the parameters in the total PDF for the fourth fit (eq. (4.20)) are fixed. Likewise, the fractions for each component  $P_4(i, q, j)$  are fixed based on the previous fits. The  $T_i$  values are fixed to those obtained by CLEO (so as to reduce the number of free parameters and improve fit behaviour). The only free parameters are  $x$ ,  $y$ ,  $\Gamma_D = 1/\tau_D$ , and the set of  $(c_i, s_i)$  values. For the latter, the information on the CLEO measurements and their uncertainties, including correlations, is incorporated as a set of correlated Gaussian constraints on the likelihood.

As in section 4.4, pseudoexperiments are used to validate the fit procedure, following all steps including the per-event decay time acceptance determination. An ensemble of 1000 experiments is generated with `cfits` [39] taking  $\Gamma_D = 2.44 \text{ ps}^{-1}$ ,  $x = -1 \times 10^{-2}$ , and  $y = +1 \times 10^{-2}$ . The mean fitted values of  $x$  and  $y$  are found to differ from the input values by  $(-0.016 \pm 0.014) \times 10^{-2}$  and  $(+0.013 \pm 0.016) \times 10^{-2}$ , respectively. The mean fitted value of  $\Gamma_D$  differs from the input value by  $(+0.0012 \pm 0.0002) \text{ ps}^{-1}$ ; although this indicates a measureable bias, it is only approximately one sixth the size of the statistical uncertainty on  $\Gamma_D$ . Since  $\Gamma_D$  is measured here only as a cross-check, this is ignored. Validation tests are also performed with a sample of pseudoevents generated with PYTHIA and EVTGEN, corresponding to approximately double the yield in data, and with a sample of events in which the full detector response was simulated with GEANT4, corresponding to approximately a quarter of the yield in data. The output is consistent with the input values of the mixing parameters supplied to the generators.

The results of the fit to data are

$$\begin{aligned} x &= (-0.86 \pm 0.53) \times 10^{-2}, \\ y &= (+0.03 \pm 0.46) \times 10^{-2}, \\ \Gamma_D &= 2.435 \pm 0.006 \text{ ps}^{-1}. \end{aligned}$$



The correlation coefficient between  $x$  and  $y$  is  $+0.37$ . The uncertainties quoted above are the statistical uncertainties estimated by the likelihood fit. They do not include any systematic effects, but they do implicitly include the propagated uncertainties on the CLEO ( $c_i, s_i$ ) parameters. These are estimated with pseudoexperiments to be in the range  $(0.05\text{--}0.15) \times 10^{-2}$ . As a check, the fit to data is repeated with the ( $c_i, s_i$ ) values fixed to those obtained by CLEO, giving  $x = (-0.73 \pm 0.48) \times 10^{-2}$  and  $y = (+0.05 \pm 0.45) \times 10^{-2}$ , with  $\Gamma_D$  unchanged. The shifts in  $x$  and  $y$  are consistent with the uncertainties associated with the CLEO parameters.

## 5 Systematic uncertainties

Further cross-checks are performed and systematic effects considered, as summarised in table 1. Several sources of systematic uncertainty are due to assumptions made for the baseline fit procedure. These uncertainties are estimated with ensembles of pseudoexperiments in which events are generated so as to mimic the effect being studied. For these tests, the systematic uncertainties on  $x$  and  $y$  are typically estimated as the sum in quadrature of the shift in the central value and the uncertainty on the shift. The fit procedure was also validated with a sample of events in which the detector response was simulated using GEANT4 as outlined in section 3; the values of  $x$  and  $y$  obtained were consistent with the input parameters.

Biases on  $x$  and  $y$  due to the fit procedure itself are assessed through the use of pseudoexperiments. The resolutions on the decay time, on the turning points, and on  $m_{12}^2$  and  $m_{13}^2$  are evaluated by generating pseudoexperiments with resolution smearing and then fitting them with the baseline procedure in which the resolution is neglected. Estimates of the resolutions are taken from data or from the full simulation based on GEANT4. The assumption that the turning point distributions of prompt and secondary signal are equivalent is tested with pseudoexperiments in which these distributions are drawn from prompt-enriched ( $\ln \chi_{\text{IP}}^2 < 1$ ) and secondary-enriched ( $\ln \chi_{\text{IP}}^2 > 3$ ) samples, respectively. The impact of neglecting variation in efficiency as a function of position in the Dalitz plot is assessed by generating pseudoexperiments with a nonuniform efficiency model, determined with full simulation, and fitting them with the baseline procedure. The efficiency is described by a polynomial function and the following variations are tested: the order of the polynomial, whether or not it is required to be symmetric about the leading diagonal in the Dalitz plot, and the use of a different event selection. The variation among models in the values of  $x$  and  $y$  is smaller than the systematic uncertainties quoted, which are based on the variation with respect to the baseline fit; in particular, the variation in  $x$  among the models is approximately  $0.01 \times 10^{-2}$ . The uncertainty associated with the model of the tracking efficiency correction  $\varepsilon(t_D)$ , discussed in section 4.3, is assessed by allowing higher-order terms in the model. Due to the absence of a  $K_s^0$  mass constraint, a small fraction of events fall outside the expected Dalitz plot boundary in the baseline procedure and an algorithm is used to assign them to a nearby bin; the effect of this is tested by instead rejecting all such events. To test the modelling of the combinatorial background,

Source	$x (\times 10^{-2})$	$y (\times 10^{-2})$
Fit bias	0.021	0.020
Decay time resolution	0.065	0.039
Turning point (TP) resolution	0.020	0.022
Invariant mass resolution	0.073	0.028
Prompt/secondary TP distributions	0.051	0.023
Efficiency over phase space	0.057	0.071
Tracking efficiency parameterisation	0.015	0.025
Kinematic boundary	0.012	0.006
Combinatorial background	0.061	0.052
Treatment of secondary $D$ decays	0.046	0.025
Uncertainty from $T_i$	0.079	0.056
Uncertainties from $(m_D, \Delta m)$ fits	0.000	0.000
Uncertainties from lifetime fit	0.020	0.043
$D^0$ background	0.001	0.006
Variation of signal components across the phase space	0.013	0.017
Total systematic uncertainty	0.171	0.134
Statistical uncertainty	0.527	0.463

**Table 1.** Systematic uncertainties on  $x$  and  $y$ . The statistical uncertainties, which include the uncertainties associated with the CLEO parameters ( $c_i, s_i$ ), are shown for comparison.

the procedure is repeated using just the data in one of the two sidebands, with the  $D^{*+}$  and  $D^{*-}$  samples separated (as in the baseline fit) or combined.

In addition, the uncertainties associated with a number of parameters that are fixed in the baseline fit are included, generally by rerunning the baseline fit repeatedly with the parameters fixed to different values obtained by smearing the nominal values randomly according to their estimated uncertainties. This procedure is used for the  $T_i$  values from CLEO, for the yield fractions estimated from the third fit to the  $(m_D, \Delta m)$  distribution, and for the decay time and  $\ln \chi_{\text{IP}}^2$  parameters fixed based on the second fit. The effects of varying the  $D^0$ - $\bar{D}^0$  composition of the prompt  $D^0$  background (via the fraction  $p_{D^0}$ ) and of using separate models of the prompt and secondary  $\ln \chi_{\text{IP}}^2$  distributions for each phase space bin are also tested.

The sum in quadrature of the systematic uncertainties is  $0.17 \times 10^{-2}$  for  $x$  and  $0.13 \times 10^{-2}$  for  $y$ .

## 6 Conclusions

The charm mixing parameters  $x$  and  $y$  have been measured using a novel method that does not require the use of an amplitude model but instead uses external measurements

of the strong phase made at an  $e^+e^-$  collider running at the  $\psi(3770)$  resonance [13]. A sample of  $pp$  collision data recorded by the LHCb experiment was used, corresponding to an integrated luminosity of  $1.0\text{ fb}^{-1}$  at a centre-of-mass energy of 7 TeV. Neglecting  $CP$  violation, the measured values are

$$\begin{aligned}x &= (-0.86 \pm 0.53 \pm 0.17) \times 10^{-2}, \\y &= (+0.03 \pm 0.46 \pm 0.13) \times 10^{-2}.\end{aligned}$$

The first uncertainties are combinations of the LHCb statistical uncertainties and those due to the CLEO measurements of the  $(c_i, s_i)$  parameters, whose effect is too small to determine precisely from the fit but is estimated to be in the range  $(0.05\text{--}0.15) \times 10^{-2}$ . The second uncertainties are systematic. The correlation coefficient between  $x$  and  $y$  for the first uncertainty is  $+0.37$ , and the systematic uncertainties are considered uncorrelated. The analysis prefers a negative value of  $x$ , but positive values are not excluded. The current HFAG world averages [5] are  $x = (+0.37 \pm 0.16) \times 10^{-2}$  and  $y = (+0.66^{+0.07}_{-0.10}) \times 10^{-2}$ .

This analysis constitutes a proof of principle that the mixing parameters can be measured in  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$  decays at LHCb without the need for an amplitude model. The statistical uncertainty will be reduced substantially by the addition of the 2012 data sample due to improvements in the software trigger, which now accepts  $D^0 \rightarrow K_S^0 \pi^+ \pi^-$  decays in which the  $K_S^0$  vertex lies outside the vertex detector, as occurs in the majority of cases. A further improvement may be obtained if charm mesons produced in semileptonic  $b$ -hadron decays are incorporated. The method does not require a detailed model of the efficiency as a function of position in the phase space, and the decay time acceptance is determined from data. Thus, the method does not rely on the extensive use of Monte Carlo simulation. This is crucial for future analyses, especially in the context of the planned LHCb upgrade where  $\mathcal{O}(10^8)$  signal events are expected [40]. To take full advantage of such a data set, more precise strong phase measurements from a charm factory running on the  $\psi(3770)$  resonance will be needed.

## Acknowledgments

We express our gratitude to our colleagues in the CERN accelerator departments for the excellent performance of the LHC. We thank the technical and administrative staff at the LHCb institutes. We acknowledge support from CERN and from the national agencies: CAPES, CNPq, FAPERJ and FINEP (Brazil); NSFC (China); CNRS/IN2P3 (France); BMBF, DFG and MPG (Germany); INFN (Italy); FOM and NWO (The Netherlands); MNiSW and NCN (Poland); MEN/IFA (Romania); MinES and FANO (Russia); MinECo (Spain); SNSF and SER (Switzerland); NASU (Ukraine); STFC (United Kingdom); NSF (U.S.A.). We acknowledge the computing resources that are provided by CERN, IN2P3 (France), KIT and DESY (Germany), INFN (Italy), SURF (The Netherlands), PIC (Spain), GridPP (United Kingdom), RRCKI (Russia), CSCS (Switzerland), IFIN-HH (Romania), CBPF (Brazil), PL-GRID (Poland) and OSC (U.S.A.). We are indebted to the communities behind the multiple open source software packages on which we depend. We

are also thankful for the computing resources and the access to software R&D tools provided by Yandex LLC (Russia). Individual groups or members have received support from AvH Foundation (Germany), EPLANET, Marie Skłodowska-Curie Actions and ERC (European Union), Conseil Général de Haute-Savoie, Labex ENIGMASS and OCEVU, Région Auvergne (France), RFBR (Russia), XuntaGal and GENCAT (Spain), The Royal Society and Royal Commission for the Exhibition of 1851 (United Kingdom).

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